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Kernelized random KISS metric learning for person re-identification



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ABSTRACT

Person re-identification is critical for human tracking in the video surveillance which has attracted more and more attention in recent years. Various recent approaches have made great progress in re-identification performance using metric learning techniques and among them, Keep It Simple and Straightforward (KISS) metric learning method has shown remarkably importance because of its simpleness and high-efficiency. The KISS method is based on an assumption that the differences between feature pairs obey the Gaussian distribution. However, for most existing features of person re-identification, the distributions of differences between feature pairs are irregular and undulant. Therefore, prior to the Guassian based metric learning step, it's important to augment the Guassian distribution of data without losing discernment. Moreover, most metric learning methods were greatly influenced by the small sample size (SSS) problem and the KISS method is no exception, which causing the inexistence of inverse of covariance matrices. To solve the above two problems, we present Kernelized Random KISS (KRKISS) metric learning method. By transforming the original features into kernelized features, the differences between feature pairs can better fit the Gaussian distribution and thus they can be more suitable for the Guassian assumption based models. To solve the inverse of covariance matrix estimation problem, we apply a random subspace ensemble method to obtain exact estimation of covariance matrix by randomly selecting and combining several different subspaces. In each subspace, the influence of SSS problem can be minimized. Experimental results on three challenging person re-identification datasets demonstrate that the KRKISS method significantly improves the KISS method and achieves better performance than most existing metric learning approaches.

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1. Introduction

Person re-identification is a sub-problem in automated video surveillance which has attracted more and more attention in the past five years because of its significance [1,2]. Given an image of one person, the objective of person re-identification is to match him/her among a large number of images of pedestrians which are captured by some different cameras. As humans, we do it all the time without paying much effort, but it's an extremely difficult task for computer because there are many factors that may influence the performance. The images of pedestrians are always low-quality, which causes the traditional biometrics, such

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http://dx.doi.org/10.1016/j.neucom.2017.08.064 0925-2312/© 2017 Elsevier B.V. All rights reserved. as face [24,38], gait [39] and iris [40], to be not available. Moreover, the appearances of pedestrians often change across camera views due to wide variations in viewpoints, illumination, and pose. The occlusions in the images also bring more difficulty to it. To deal with this problem, researchers have proposed many different methods in which the most common research direction is feature representation. It aims to extract robust and effective features from each pedestrian image. The features can be hand-crafted [4–6,18–20,29] or learned [3,17,28,30–34,44,59,60] based on visual information like color, texture, and shape. It has been proved that among all the visual information, color plays the most important role for person re-identification.

After extracting distinguishing features, the next step is naturally to compute the distances between different feature pairs and choose the smallest one as the matching image pair. Different distances, such as Euclidean distance, cosine distance, and Mahalanobis distance, are usually used for pedestrian matching. The



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procedure of learning a discriminant distance function is called metric learning. Researchers have proposed many metric learning algorithms for person re-identification which can be summarized to several groups according to their basic theories. For instance, based on the theory of Fisher Discriminant Analysis (FDA), some metric learning methods such as, Local Fisher Discriminant Analysis (LFDA) [7], Marginal Fisher Analysis (MFA) [8] and Cross-view Quadratic Discriminant Analysis (XQDA) [6], were proposed. Most of them tried to project descriptors into a more discriminatory subspace based on an eigenvalue resolution. By adding special constraints to the distance functions, some methods like Pairwise Constrained Component Analysis (PCCA) [9], Large Margin Nearest Neighbor (LMNN) [10] and Locally-Adaptive Decision Functions (LADF) [11] were proposed. Different constraints in the methods show different performances when they are applied for person re-identification. Some kernel-based metric learning methods also have been proposed to improve the identification ability of features because of the non-linearity in the appearance of pedestrians [5,8,21,35,41-43]. The benefits of kernel-based methods for person re-identification are twofold: (1) True matches and wrong matches become more separable in the nonlinear space; (2) the kernel can be chosen flexibly. In addition, value difference metric (VDM) [49] and its varieties [50-58] have been proposed to find reasonable distance between each pair of instances with nominal attributes only.

It is specially worth mentioning that based on the maximum likelihood (ML) estimation, Keep It Simple and Straightforward (KISS) metric learning method [12] was proposed. The KISS metric learning method is both efficient and effective by considering a log likelihood ratio test of two Gaussian distributions as the distance between a feature pair. With a simplified and efficient solution, the covariance matrix of Mahalanobis distance is obtained by computing the difference between the inverse of two covariance matrices. Although the KISS metric learning method has been widely applied to the person re-identification because of its simpleness and effectivenes36s, there are still two problems. Firstly, there is an assumption of the KISS metric learning method that pairwise differences are agreeing with Gaussian distribution, if not, the performance would decrease. Second, most metric learning methods suffer from SSS (small sample size) problem because the size of samples is always much smaller than the dimensionality of features, and the KISS metric learning method is no exception. In the training procedure of the KISS method, the covariance matrices may be singular when the training size of data is small, resulting in the inexistence of the inverse of covariance matrices. How to augment the Guassian distribution of data without losing discernment and how to accurately estimate the inverse of covariance matrix in the case of small samples are still open problems which exsit in various fields. To overcome the inverse of covariance matrix estimation problem, unsupervised dimensionality reduction is required as the most common method. In [12], before applying the KISS metric learning method, the PCA (Principal Component Analysis) was used to project the concatenated descriptors into a 34-dimensional subspace which was greatly less than the original feature dimension. However, in the procedure of dimension reduction, the abandon dimensionalities also have the ability of distinguishing pedestrians, and thus the unsupervised dimensionality reduction will make the learned distance less discriminative. There are some improved metric learning methods based on the KISS method such as Regularized Smoothing KISS (RS-KISS) [13], minimum classification error-KISS (MCE-KISS) [14], Regularized Bayesian Metric Learning (RBML) [15] and Dual-Regularized KISS (DR-KISS) [36]. Although improvement of performance can be seen, the problems mentioned above are still unsolved.

In this paper, based on the KISS method we propose a novel metric learning method called Kernelized Random KISS (KRKISS) in which a special kernel technique is used to augment the Gaussian distribution f data by transforming the original features into kernelized features, and a random subspace ensemble method is applied tosolve the inverse of covariance matrix estimation problem in the case of small samples. It contains the advantages of the KISS metric learning method and the kernel theory, and avoids using dimensionality reduction techniques or regularization techniques which will cause a loss to discernment of features. We have introduced above that the KISS method was proposed under the assumption of Gaussian distribution. When the original features are transformed into kernelized features by our proposed method, they can better fit Gaussian distribution and obtain more discrimination for person re-identification. It is the inspiration and motivation that why we utilize the transformation to improve the KISS metric learning method. As far as we know, it is the first of its kind to improves the KISS method by augmenting the Guassian model of data. Moreover, the covariance matrix usually be singular when the size of samples is small, which would cause the inexistence of the inverse of covariance matrix. The random subspace ensemble method we applied to obtain exact estimation of covariance matrix is simple but very efficient, which decreases the feature dimension by randomly selecting and combining several different subspaces. In each subspace, the influence of SSS problem can be minimized. It also works to other covariance matrix based metric learning methods. Experiments with five kinds of features (LOMO [6], kCCA [5], SCNCD [4], ELF18 [19] and FFN [17]) on different datasets show that our proposed KRKISS metric learning algorithm can improve the identification accuracy obviously and get better performance than other state-of-the-art metric learning methods in most of experiments.

The main contributions of our work are summarized as below:

- (1) We augment the Gaussian distribution of data by transforming the original feature into special kernelized feature. By doing so, the differences between feature pairs can better fit the Gaussian distribution, which is more aligned with the metric learning methods based on Gaussian distribution assumption.
- (2) We apply a random subspace ensemble method to take the place of regularization methods in the procedure of estimating the inverse of covariance matrix. In each randomly selected subspace, the influence of SSS problem can be minimized and the inverse of covariance matrix can be estimated more accurately.

The rest of the paper is organized as follows. Section 2 introduces some related works about person re-identification. Section 3 proposes our Kernelized Random KISS (KRKISS) metric learning method. Experiments on three public datasets are described in Section 4. Finally, the conclusion is draw in Section 5.

2. Related work

The proposed methods for person re-identification can be generally categorized into two groups which have been briefly described in Section 1. The first group of methods focuses on designing features which are robust to viewpoint change, illumination change and other variations in pedestrian's images. There are many useful features like the ensemble of localized features (ELF) [3], salient color names based color descriptor (SCNCD) [4], Local Maximal Occurrence (LOMO) [6] and hierarchal Gaussian descriptor (GOG) [29] which have been proposed and applied in person re-identification successfully. However, they are still not robust enough to huge variations in pedestrian's images. Therefore, more and more researchers turn attention to the second group of methods (metric learning) which focuses on learning a robust distance or similarity function to deal with the matching problem.

Many effective metric learning algorithms have been proposed and widely applied to solve the person re-identification problem. Davis et al. [37] proposed information-theoretic metric learning (ITML) from the perspective of information theoretic. Weinberger et al. [10] proposed the large-margin nearest neighbor metric (LMNN) to improve the performance of the kNN classification. In [9], an algorithm for learning distance metric called Pairwise Constrained Component Analysis (PCCA) was proposed. It projected the features into a low-dimensional space where the distance between pairs of data points respects the desired constraints. In addition, some kernel based metric learning methods have been proposed in recent years. Xiong et al. [8] proposed kernel LFDA metric learning method which performs better than LFDA [7]. Lisanti et al. [5] applied a learning technique based on Kernel Canonical Correlation Analysis (KCCA) which finds a common subspace between the proposed descriptors extracted from disjoint cameras and then projects them into a new descriptors space. Hu et al. [48] proposed a new deep transfer metric learning (DTML) method to learn a set of hierarchical nonlinear transformations by transferring discriminative knowledge from the labeled source domain to the unlabeled target domain. Duan et al. [47] proposed a deep localized metric learning (DLML) approach by learning multiple fine-grained deep localized metrics. In [21], kernel technique was utilized to learning a kernelized discriminative null space.

Moreover, Köstinger et al. [12] proposed the KISS metric learning algorithm which considered a log likelihood ratio test of two Gaussian distributions and obtained a simplified and efficient solution to solve it. There are also some improved methods based on the KISS method. In [13], a regularized smoothing KISS metric learning (RS-KISS) was presented which seamlessly integrated smoothing and regularization techniques to estimate covariance matrices. In [14], the KISS metric learning method was improved based on the minimum classification error (MCE). Tao et al. [36] proposed a Dual-Regularized KISS (DR-KISS) metric learning by regularizing the two covariance matrices and theoretically demonstrated that the proposed regularization method is robust for generalization. However, the above KISS based improved methods focus on solving the SSS problem. They may still perform poorly when the features do not obey the Gaussian Distribution.

In this paper, we focus on the second group of methods, i.e. metric learning. Aiming at augmenting Guassian distribution of data and exactly estimating the inverse of covariance matrix, we propose Kernelized Random KISS (KRKISS) metric learning method.

3. Proposed approach

3.1. Kernelizd random KISS metric learning (KRKISS)

The person re-identification problem can be formulated as follows. Let $(x_i, y_i)_{i=1}^m$ denote all training samples, where x_i is the *i*-th sample, y_i is the identity of the corresponding person and *m* is the number of training samples. The goal of metric learning methods is to learn a generic distance function $d(\cdot)$ from the training samples that outputs smaller value for samples of the same person and larger value for different people. During the test phase, given a probe pedestrian image and a set of gallery images, they are transformed into descriptors firstly. Then we use the distance function $d(\cdot)$ to measure distance from all of them, and rank the gallery images according to their distance to the probe image. Usually we use the Mahalanobis distance function to measure the distance between two samples x_i and x_i , such as

$$d(x_i, x_j) = (x_i - x_j)^T M(x_i - x_j)$$

thus we need to learn a robust and discriminative matrix M from training samples. In the KISS metric learning method [12], under a assumption that the differences between features pairs obey



Fig. 1. The typical distribution of differences between feature pairs. The blue bar represents the actual distribution and the red line represents the computed Gaussian distribution. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

Guassian distribution, the matrix *M* was calculated by

$$M = \Sigma_1^{-1} - \Sigma_0^{-1}$$

where Σ_1 is the covariance matrix of differences between same person and Σ_0 is the covariance matrix of differences between different people.

It is an simple and efficient metric learning algorithm, however, the commonly used features for person re-identification do not obet the assumption that the differences between feature pair are Gaussian distributed. For example, the typical distribution of differences between feature pairs are shown in Fig. 1. As we can see, the actual distribution is at odds with the computed Gaussian distribution, which causes a significant decrease in performance of Guassian distribution based metric learning methods. The Guassian distribution assumption puts forward high demand to the feature representation. Moreover, the sample size is small in most cases, and thus the estimation of inverse of covariance matrix is inaccurate, or not exist. Some regularization methods have been proposed to solve it [13,14,15,36]. However, the existing regularization methods are still not robust enough because the weights of regularization are hand-crafted. To solve those two problems, we propose Kernelized Random KISS (KRKISS) metric learning method. For convenience, Table 1 lists frequently used notations and descriptions in the paper.

The first procedure of KRKISS is to transform the original features into kernelized features. Given a function $\Phi(x_i)$ which maps the sample x_i into an unknown space. Different with the common usage of kernel trick, we utilize the inner products of $\Phi(x_i)$ and a training sample feature $\Phi(x_i)$ as a novel feature of sample x_i . For example, $\Phi(x_i)^T \Phi(x_i)$ denotes the *j*-th novel feature, and thus the sample x_i can be transformed into $[\Phi(x_i)^T \Phi(x_1), \Phi(x_i)^T \Phi(x_2), \dots, \Phi(x_i)^T \Phi(x_m)]$ when there are *m* training samples. Given a kernel $k(x_i, x_j) = \langle \Phi(x_i), \Phi(x_j) \rangle = \Phi(x_i)^T \Phi(x_j),$ function xi can be transformed into kernelized features $\tilde{x}_i = \mathbf{K}(x_i, \mathbf{X}) =$ $[k(x_i, x_1), k(x_i, x_2), \dots, k(x_i, x_m)]^T \in \mathbb{R}^{m \times 1}$. In the same way, the training samples X and the testing samples Z can be transformed into $\mathbf{\tilde{X}} = [\mathbf{K}(x_1, \mathbf{X}), \mathbf{K}(x_2, \mathbf{X}), \dots, \mathbf{K}(x_m, \mathbf{X})] \in \mathbf{R}^{m \times m}$ and $\mathbf{\tilde{Z}} = [\mathbf{K}(z_1, \mathbf{X}), \mathbf{K}(z_2, \mathbf{X}), \dots, \mathbf{K}(z_n, \mathbf{X})] \in \mathbf{R}^{m \times n}$ respectively. Unless stated, otherwise, RBF kernel is used in our method with kernel width determined automatically using the mean pairwise distance of samples. After transforming the original features into kernelized features, the actual distribution of differences between kernelized feature pairs better fit the computed Gaussian distribution.

Table 1						
Frequently used r	notations	and	descriptions	in	the	paper.

Notation	Description	Notation	Description
\mathbf{X}/x_i m	Training samples Number of training samples	Z /z _i n	Testing samples Number of testing samples
$egin{array}{lll} {f ilde X}/{ ilde x}_i \ { ilde x}_{ij} \end{array}$	Kernelized features of training samples Difference of kernelized feature pair and \tilde{x}_j	$oldsymbol{ ilde{Z}}/\widetilde{z}_i \ \widetilde{z}_{ij}$	Kernelized features of testing samples Difference of kernelized feature pair \tilde{z}_i and \tilde{z}_j

Moreover, there are many advantages of kernelized features which will be described in detail and analyzed in next section.

Given a pair of testing samples z_i and z_j that we need to decide if they are the same person. They can be transformed to kernelized features \tilde{z}_i and \tilde{z}_j according to the above method. Let H_0 denotes the hypothesis that the testing feature pair (\tilde{z}_i and \tilde{z}_j) is dissimilar (\tilde{z}_i and \tilde{z}_j are different people), H_1 denotes the hypothesis that the feature pair (\tilde{z}_i and \tilde{z}_j) is similar (\tilde{z}_i and \tilde{z}_j are the same people). If \tilde{z}_{ij} satisfies the Gaussian distribution, we have

$$p(\tilde{z}_{ij}|H_k) = \frac{1}{(2\pi)^{m/2} |\Sigma_k|^{1/2}} \exp\left(-\frac{1}{2}\tilde{z}_{ij}^T \Sigma_k^{-1} \tilde{z}_{ij}\right)$$
(1)

where $k \in \{0, 1\}$, $p(\tilde{z}_{ij}|H_0)$ and $p(\tilde{z}_{ij}|H_1)$ denotes the probability of \tilde{z}_{ij} under the hypothesis of H_0 and H_1 respectively, Σ_0 is the covariance matrix of interpersonal differences and Σ_1 is the covariance matrix of intrapersonal differences.

The two covariance matrices can be estimated by the training samples. Define y_{ij} as the indicated variable of two training samples \tilde{x}_i and \tilde{x}_j : $y_{ij} = 1$ if \tilde{x}_i and \tilde{x}_j are the same person and $y_{ij} = 0$, otherwise. So we have

$$\Sigma_{0} = \frac{1}{N_{0}} \sum_{y_{ij}=0} \tilde{x}_{ij} \tilde{x}_{ij}^{T} = \frac{1}{N_{0}} \sum_{y_{ij}=0} (\tilde{x}_{i} - \tilde{x}_{j}) (\tilde{x}_{i} - \tilde{x}_{j})^{T}$$

$$\Sigma_{1} = \frac{1}{N_{1}} \sum_{y_{ij}=1} \tilde{x}_{ij} \tilde{x}_{ij}^{T} = \frac{1}{N_{1}} \sum_{y_{ij}=1} (\tilde{x}_{i} - \tilde{x}_{j}) (\tilde{x}_{i} - \tilde{x}_{j})^{T}$$
(2)

where N_0 denotes the number of interpersonal differences, N_1 denotes the number of intrapersonal differences.

In order to decide if \tilde{z}_i and \tilde{z}_j are the same person, the logarithm of the ratio between two posterior probabilities is defined as follows:

$$\delta(\tilde{z}_{ij}) = \log\left(\frac{p(H_0|\tilde{z}_{ij})}{p(H_1|\tilde{z}_{ij})}\right)$$
(3)

According to the Bayes formula, (3) can be transformed to

$$\delta(\tilde{z}_{ij}) = \log\left(\frac{p(\tilde{z}_{ij}|H_0)}{p(\tilde{z}_{ij}|H_1)}\right) + \log\left(\frac{p(H_0)}{p(H_1)}\right)$$
(4)

Given (1), (4) can be rewritten as

$$\delta(\tilde{z}_{ij}) = \frac{1}{2} \tilde{z}_{ij}^{T} (\Sigma_{1}^{-1} - \Sigma_{0}^{-1}) \tilde{z}_{ij} + \frac{1}{2} \log\left(\frac{|\Sigma_{1}|}{|\Sigma_{0}|}\right) + \log\left(\frac{p(H_{0})}{p(H_{1})}\right)$$
(5)

So \tilde{z}_i and \tilde{z}_j are the same people while the value of $\delta(\tilde{z}_{ij})$ is negative and a positive value of $\delta(\tilde{z}_{ij})$ indicates \tilde{z}_i and \tilde{z}_j are different people. By dropping the constant in (5), we have

$$\delta(\tilde{z}_{ij}) = \tilde{z}_{ij}^T (\Sigma_1^{-1} - \Sigma_0^{-1}) \tilde{z}_{ij}$$
(6)

The smaller the value of $\delta(\tilde{z}_{ij})$ is, the bigger the probability of \tilde{z}_i and \tilde{z}_j belong to same person. It is similar with the Mahalanobis distance in form, and thus the distance between two testing samples z_i and z_j is

$$d(z_i, z_j) = (\tilde{z}_i - \tilde{z}_j)^T (\Sigma_1^{-1} - \Sigma_0^{-1}) (\tilde{z}_i - \tilde{z}_j)$$
(7)

The dimension of kernelized feature is the same as the number of training samples, which is greatly less than the dimension of original feature in most of scenes. However, the SSS problem still remains, resulting in the inexistence of the inverse matrices Σ_0^{-1} and Σ_1^{-1} . The common solution is to add a regularization process for covariance matrix such as

$$\Sigma_{0} = (1 - \alpha_{0})\Sigma_{0} + \alpha_{0} \frac{\operatorname{trace}(\Sigma_{0})}{d}I$$

$$\Sigma_{1} = (1 - \alpha_{1})\Sigma_{1} + \alpha_{1} \frac{\operatorname{trace}(\Sigma_{1})}{d}I$$
(8)

where α_0 and α_1 denotes the weights of regularization. The significance of this regularization method has been demonstrated theoretically in [36]. However it is not robust enough because the parameters α_0 and α_1 are hand-controlled.

To address this problem, we propose a random subspace ensemble method which projects the training features into different randomly selected subspaces. Then the inverse matrices are computed by all the training samples in each subspace. When the dimension of subspace is greatly less than the number of training samples, we can obtain the inverse matrices without the SSS problem. The final distance is computed by

$$d(z_i, z_j) = \sum_{k=1}^{L} \left(D_k(\tilde{z}_i) - D_k(\tilde{z}_j) \right)^T M_k(D_k(\tilde{z}_i) - D_k(\tilde{z}_j))$$
(9)

$$M_{k} = \Sigma_{1}^{-1} - \Sigma_{0}^{-1}$$

= $\frac{1}{N_{1}} \sum_{y_{ij}=1} D_{k}(\tilde{x}_{ij}) D_{k}(\tilde{x}_{ij})^{T} - \frac{1}{N_{0}} \sum_{y_{ij}=0} D_{k}(\tilde{x}_{ij}) D_{k}(\tilde{x}_{ij})^{T}$ (10)

where *L* denotes the number of selected subspaces and D_k denotes a randomly selected subspace. The diagram of KRKISS metric learning method is shown in Fig 2 and the complete KRKISS algorithm is summarized in Algorithm 1.

3.2. Discussion of kernelized random KISS method

The Kernelized Random KISS (KRKISS) method has the main advantages as below:

(1) When the original features are transformed into kernelized features, the differences between feature pairs can better fit Gaussian distribution.

For example, given the LOMO feature $[d_1, d_2, \dots, d_{26960}]^T =$ $X \in R^{26960 \times 632}$ extracted from VIPeR dataset [25], where 26,960 is the dimensionality of the LOMO feature, 632 is the number of samples and $d_i \in \mathbb{R}^{1 \times 632}$ denotes the *j*-th dimension data. For every one-dimensional data d_k , it is able to calculate two sets of differences dif_0 and dif_1 according to the labels of samples, where dif₀ denotes a set of differences between interperson feature pairs and dif_1 denotes a set of differences between intraperson feature pairs. Since the means of dif_0 and dif_1 are both zero, we can obtain two Gaussian distributions by calculating the variances of dif_0 and *dif*₁ respectively. Comparing the concrete probability distributions of differences with the computed Gaussian distribution, we can consider the set of differences obeys the Gaussian distribution if they are similar in shape. The typical probability distributions of four original features (LOMO [6], kCCA [5], SCNCD [4] and ELF18 [19]) are shown in Fig 3a and b. The blue bars are concrete probability distributions of differences and the red line is the computed



Fig. 2. The diagram of KRKISS metric learning method.

Algorithm 1

Kernelized Random KISS (KRKISS).

Input: Training samples $(x_i, y_i)_{i=1}^m$, testing samples $(z_i)_{i=1}^n$, the number of se lected subspaces L and the dimension of subspace dOutput: Some Mahalanobis matrices M_k , (k = 1, 2, ..., L) and the distance be tween two test samples z_i and z_j 1: while i < m2: $i \leftarrow i + 1$; 3: Calculate the kernelized feature of *i*th training sample by $\tilde{x}_i = \mathbf{K}(x_i, \mathbf{X}) = [k(x_i, x_1), k(x_i, x_2), ..., k(x_i, x_m)]^T$; 4: end while 5: while t < n6: $t \leftarrow t + 1$; 7: Calculate the kernelized feature of *t*-th testing sample by $\tilde{z}_t = \mathbf{K}(z_t, \mathbf{X}) = [k(z_t, x_1), k(z_t, x_2), ..., k(z_t, x_m)]^T$; 8: end while 9: while k < L10: Randomly select a *d*-dimension subspace D_k from kernealized feature space; 11: Calculate the *k*-th Mahalanobis matrix M_k according to Eq. (10); 12: end while

13: Calculate the distance between two test samples z_i and z_j according to Eq. (9)

Table 2	
The Lilliefors test results of two features.	
Features /Lilliofers test LOMO	1-004

Table 3The variance ratio of two features.

Features /Lilliefors test	LOMO kCCA		Features	LOMO		kCCA			
	Original features	Kernelized features	Original features	Kernelized features		Original features	Kernelized features	Original features	Kernelized features
Lilliefors test of dif_1 Lilliefors test of dif_0	0.0462 0.0355	0.3497 0.3956	0.4673 0.4691	0.6139 0.8149	variance ratio $\frac{\sigma_0}{\sigma_1}$	1.1502	1.4389	1.2429	2.4177

Gaussian distribution. As we can see in Fig 3a and b, the probability distributions of the differences between original feature pairs are irregular and most of them are concentrated around the zero value. The computed Gaussian distributions do not fit the probability distributions of the differences since the values of variance are higher when the distributions are irregular. In the same way, it is able to calculate the differences between kernelized feature pairs and the typical probability distributions of four kernelized features (LOMO, kCCA, SCNCD, ELF18), as shown in Fig 3c and d. When the original features are transformed into kernelized feature, the probability distributions of the differences will be more regular and better fit to the computed Gaussian distribution in shape.

Moreover, we have applied the Lilliefors test [45] to test the null hypothesis that the data come from a normally distributed population. The Lilliefors test is a normality test based on the Kolmogorov-Smirnov test with mean and variance unknown. We set the value of $\frac{n_0}{N}$ to represent the degree of distribution, where n_0 denotes the number of one-dimensional feature which is consistent with the outcome of Lilliefors test, N denotes the dimensionality of features. The value of $\frac{n_0}{N}$ is larger and the features better fit to Gaussian distribution. The results are shown in Table 2. As we can see in Table 2, the value of $\frac{n_0}{N}$ increases significantly when the original features are transformed into kernelized feature.

It demonstrates that the kernelized feature better fit to the Gaussian distribution and is more suitable for the Guassian distribution assumption based metric methods.

(2). The kernelized features always have higher recognition capability in the non-linear space.

The variance ratio of two differences sets dif_0 and dif_1 can be seen as a performance index of discernment [6]. The distributions of dif_0 and dif_1 are shown in Fig 4. σ_0 denotes the variance of dif_0 and σ_1 denotes the variance of dif_1 . The higher the variance ratio $\frac{\sigma_0}{\sigma_1}$ is, the more discriminative the feature is. The value of variance ratio for every one-dimensional feature was calculated and the average value was taken. The results are shown in Table 3. Since the mean variance ratio of kernelized feature is higher than the original feature, we consider that when the original features are transformed into the kernelized features, they would become more distinguishable.

(3) The robustness to the variation of parameters.

There are two parameters in KRKISS method, the number of selected subspaces *L* and the dimension of subspace *d*. Experientially we set $d = \frac{m}{4}$ to guarantee the existence of the inverse of covariance matrices in subspace. Next, we conduct some experiments with different parameters to compare the influence of *L* in KRKISS method. Fig 5 shows the experimental results of LOMO feature and kCCA feature respectively. The blue line represents the impact



Fig. 3. The probability distributions. The blue bars are concrete probability distributions of differences and the red line is the computed Gaussian distribution of differences. (a) Differences of original features between interperson image pairs. (b) Differences of original features between interperson image pairs. (c) Differences of kernelized features between interperson image pairs. (d) Differences of kernelized features between interperson image pairs. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

of subspaces number on rank-1 matching rate and the red line represents the impact of regularized weight on rank-1 matching rate.

As we can see in Fig 5, the matching rate of regularized method has obvious fluctuations which cause the difficulty to choose a better parameter. Instead, the matching rate of random subspace ensemble method always achieves near the maximum when the number of subspaces increases,. Taking into account of the efficiency of algorithm, we set $L = \frac{d}{2}$, and thus all the parameters in the KRKISS method need not to be adjusted manually.

(4) Low computational complexity.

Suppose the *m* training samples consist of *a* images captured from one view and *b* images captured from the other view (m = a + b). In KRKISS method, the computation of feature transform requires $O(dm^2)$ multiplication operations and for each randomly selected subspace, the computation of the two covariance matrices Σ_I and Σ_E require $O(Nkm^2)$ and $O(abm^2)$ multiplication operations, respectivesly, where $N = \max(a, b)$, *d* denotes the dimension of original feature and *k* represents the average number of images in each person. Therefore, the total computation of the two covariance matrices Σ_I and Σ_E in KRKISS method require $O(LNkm^2)$ and $O(Labm^2)$ multiplication operations, respectivesly.



Because Σ_I and Σ_E can be computed directly from sample mean and covariance of each class and all classes according to the computing method in [6], the total computations of Σ_I and Σ_E can be both reduced to $O(LNm^2)$. Adding the computation of feature transform, the computation of KRKISS requires $O(LNm^2 + dm^2)$ multiplication operations totally. Compared with the computation of KISS method ($O(Nd^2)$), the computation of KRKISS is even less when the dimension of original feature is much larger than the size of training samples. It is a common case in the applications of person reidentification and for example, in [6], the sizes of training samples in the expriments of VIPeR and QMUL dataset are 632 and 250, respectively, while the dimension of feature is 26,960 and much larger than the size of training samples. The efficientness of KRKISS is also validated through a related experiment in next section.

4. Experiments

In this section, we conduct experiments to evaluate the KRKISS method and compare it with the state of the art methods. All experiments were tested on publicly and standard datasets, namely VIPeR [25], PRID 450S [26] and CUHK01 [27]. For all the



Fig. 6. Example images from VIPeR dataset [25].

experiments, we used the cumulative matching characteristic (CMC) curve to evaluate the identification performance which is the most widely used evaluation methodology for person reidentification. It shows how performance improves as the number of requested images increases.

The features used in our experiments includes LOMO [6], kCCA [5], FNN [17], SCNCD [4], ELF18 [19]. Moreover, we propose a novel feature which consists of different color, texture and shape feature. The extracted method of color feature is same as [5] and [29]. Then we extract the Scale Invariant Local Ternary Pattern (SILTP) [6] description as the texture feature. The shape feature is Histogram of Oriented Gradient (HOG) descriptor [23] and the extracting procedure is similar to [22]. We use the novel feature as original features of pedestrians in the following experiments unless noted otherwise.

4.1. Experiments on the VIPeR dataset

VIPeR [25] is a challenging person re-identification dataset which has been widely used for performance evaluation. It contains 632 pedestrian image pairs which are captured from two different camera views. The cameras have different viewpoints and illumination variations exist between them. All images have been scaled to 128×48 pixels. Fig 6 shows some example images from this dataset. In the experiments, the VIPeR dataset was randomly divided into half for training and the other half for testing. In



Fig. 5. CMC curves on the VIPeR dataset (P = 316). (a) Comparison of different parameters by using the LOMO feature. (b) Comparison of different parameters by using the kCCA feature.



Fig. 7. CMC curves on the VIPeR dataset (P = 316). (a) Comparison of KISS based metric learning methods by using the LOMO feature. (b) Comparison of KISS based metric learning methods by using the kCCA feature.

Comparison of different metric learning methods on the VIPeR database (P = 316). The cumulative matching scores (%) at rank 1, 5, and 10 are listed. Red and blue numbers are the best and second best results, respectively.

Feature/Method	LOMO(d	= 26,960)		kCCA(d	kCCA(d = 5138)		SCNCD(a	SCNCD(d = 70)			ELF18(d = 8064)		
	r = 1	<i>r</i> = 5	<i>r</i> = 10	<i>r</i> = 1	<i>r</i> = 5	<i>r</i> = 10	r = 1	<i>r</i> = 5	<i>r</i> = 10	r = 1	<i>r</i> = 5	<i>r</i> = 10	
KRKISS	41.7	73.7	85.1	46.0	76.0	87.8	41.3	73.1	85.0	40.3	72.7	84.8	Ours
NFST [21]	40.1	70.3	82.3	44.9	74.5	86.0	36.7	67.6	80.0	37.8	70.3	84.0	2016 CVPR
MLAPG [16]	39.5	70.0	82.4	32.3	61.8	75.8	25.8	57.6	73.1	18.0	44.7	61.5	2015 ICCV
XQDA [6]	39.0	69.3	81.3	33.5	62.3	74.4	24.3	50.9	65.2	30.7	59.2	72.9	2015 CVPR
MFA [8]	35.4	67.9	81.0	40.4	70.1	83.1	37.9	68.9	81.4	36.8	70.3	83.0	2014 ECCV
kLFDA [8]	36.3	65.1	77.1	39.2	67.7	79.8	36.9	68.5	81.1	32.6	60.7	71.7	2014 ECCV
RBML [15]	33.4	65.1	78.6	37.7	67.2	79.5	37.7	65.2	77.6	33.9	63.8	77.1	2014 ECCV
LFDA [7]	37.6	68.7	81.3	34.8	64.6	77.5	16.3	29.8	36.6	33.1	63.3	76.8	2013 CVPR
RS-KISS [13]	33.0	66.7	80.4	36.2	67.0	79.8	37.1	67.2	79.7	34.2	63.6	76.6	2013 IEEE
KISSME [12]	32.3	64.6	77.9	35.8	65.8	78.8	38.7	69.8	82.8	32.7	63.5	78.5	2012 CVPR

order to obtain a precise experimental result, this procedure was repeated 10 times and an average value was taken.

4.1.1. Comparison of metric learning algorithms

Firstly, in order to demonstrate the improvement to KISS method, we compare our KRKISS method with three KISS based methods (KISSME [12], RS-KISS [13] and RBML [15]) by using two features (LOMO [6] and the kCCA [5]) respectively. Both features have greatly identification ability for person re-identification. The experimental results are shown in Fig 7. In the experiment of the LOMO feature, it can be seen that our proposed method (KRKISS) is obviously better than other KISS based methods, achieving a rate of 41.7% at rank = 1. In the experiments of the kCCA feature, our proposed KRKISS method improves the rank-1 accuracy from 37.7% to 46.0%. Both experimental results show the greatly improvement of KRKISS method to KISS based methods and demonstrate that the kernelized feature transform and the random subspace ensemble method are necessity for KISS based methods.

Next we compare the KRKISS method with some state-of-theart metric learning methods which were reported having good performance on the VIPeR dataset. The compared metric learning methods includes NFST [21], MLAPG [16], XQDA [6], MFA [8], kLFDA [8], RBML [15], LFDA [7], RS-KISS [13] and KISSME [12]. For testing the robustness of metric learning methods to different features, we apply four kinds of features (LOMO [6], kCCA [5], SCNCD [4] and ELF18 [19]) for all metric learning methods in the experiments. The experimental results are shown in Table 4 and Fig 8. It can be seen that our proposed KRKISS method achieves the best matching rates in all experiments with four features, which demonstrates that the KRKISS method not noly

Table 5				
Fraining time	(seconds) of a	metric lear	ning algor	ithms.

	KRKISS	NFST	MLAPG	XQDA	MFA	kLFDA	KISSME	LFDA
Time	5.04	2.48	40.9	3.86	2.58	2.74	7.41	229.3

has better performance than all the compared methods, but also has good stability and strong robustness to different kinds of features. Meanwhile, the average training time comparison of metric learning methods is shown in Table 5 (including subspace learning time). The training was performed on a notebook PC with an Intel i5-3210 @2.50 GHz CPU. Table 5 shows that the KRKISS is efficient and the time consumption is even less than KISSME [12] method. The theoretical analysis and the experimental results both show the efficientness of our proposed KRKISS method. Although it needs to compute several Mahalanobis matrices in different subspaces, the KRKISS method is still efficient because the dimension of subspace is greatly less than original features.

4.1.2. Comparison to the state of the art

In order to demonstrate that our proposed method (KRKISS) can achieve state-of-the-art performance for person re-identification, we compare it with other state-of-the-art methods like GOG [29], FNN [17], NFST [21], etc. The compared methods include two feature represent based methods [6,29] and five metric learning based methods [6,16,21]. In the experiment, all the parameter settings, the used features and the metric learning method were refer to their own paper. The experimental result is shown in Table 6 and Fig 9. It can be seen that our proposed method (KRKISS) achieves the best performance while it improves the most



Fig. 8. CMC curves on the VIPeR dataset (P = 316). (a) Comparison of different metric learning methods by using the LOMO feature. (b) Comparison of different metric learning methods by using the kCCA feature. (c) Comparison of different metric learning methods by using the ELF18 feature.

Comparison of state-of-the-art results reported with the VIPeR database (P = 316). The cumulative matching scores (%) at rank 1, 5, 10, and 20 are listed. The best and the second best scores are respectively shown in red and blue.

Method	r = 1	<i>r</i> = 5	<i>r</i> = 10	<i>r</i> = 20	Reference
KRKISS	54.2	83.5	91.3	96.4	Ours
GOG [29]	47.0	75.1	85.4	93.9	2016 CVPR
NFST [21]	42.3	71.5	82.9	92.1	2016 CVPR
MLAPG [16]	39.5	70.0	82.4	92.8	2015 ICCV
LOMO + XQDA [6]	39.0	69.3	81.3	91.6	2015 CVPR
RBML [15]	33.4	65.1	78.6	90.2	2014 ECCV
RS-KISS [13]	33.0	66.7	80.4	91.0	2013 IEEE
KISSME [12]	32.3	64.6	77.9	89.5	2012 CVPR

important index for person re-identification (rank-1 accuracy) from 47.0% to 54.2%. It proves our model can effectively improve the performance of person *Re*-ID.



Fig. 9. The CMC curves and rank-1 matching rates on the VIPeR dataset.

4.2. Experiments on the PRID 450S dataset

In this section, we conduct some experiments to evaluate the KRKISS method on the PRID 450S dataset [26]. The PRID 450S dataset consists of 450 images pairs of pedestrians with significant differences in background, viewpoint and illumination. It has been recently released and is regarded as more realistic. Some example images from the PRID 450S dataset are shown in Fig 10. In the experiments, the PRID 450S dataset was randomly divided into half for training and half for testing and the procedure was repeated 10 times to record an average performance.

4.2.1. Comparison of metric learning algorithms

Firstly, we compare our proposed KRKISS method with other three KISS based methods (KISSME [12], RS-KISS [13] and RBML [15]). The used features include two efficient features, LOMO [6] and FFN [17]. Among them, the LOMO feature is artificially designed and the FNN feature is learned by a deep learning method. The experimental results are shown in Fig 11. As we can see in Fig 11, whether using the LOMO feature or the FNN feature, the red line which shows the cumulative matching rate curve of KRKISS method is higher than other lines. It demonstrates

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Table 7

Comparison of different metric learning methods on the PRID 450S database (P = 225). The cumulative matching scores (%) at rank 1, 5 and 10 are listed. The best and the second best scores are respectively shown in red and blue.

Method/Feature	LOMO(d	= 26,960)		FFN(d =	FFN(d = 4096)		SCNCD(a	SCNCD(d = 70)			ELF18(d = 8064)		
	r = 1	<i>r</i> = 5	<i>r</i> = 10	<i>r</i> = 1	<i>r</i> = 5	<i>r</i> = 10	r = 1	<i>r</i> = 5	<i>r</i> = 10	r = 1	<i>r</i> = 5	<i>r</i> = 10	Reference
KRKISS	58.0	83.8	90.9	53.0	77.5	85.8	43.7	71.0	81.6	31.9	59.5	72.3	Ours
NFST [21]	57.1	82.8	89.8	54.4	79.0	86.9	32.0	57.7	68.6	29.5	54.1	67.4	2016 CVPR
MLAPG [16]	55.5	80.1	88.3	22.7	46.5	57.9	30.4	60.3	73.6	8.8	23.8	35.2	2015 ICCV
XQDA [6]	55.9	80.6	88.0	48.4	71.6	80.9	16.1	40.7	53.6	27.8	51.5	63.0	2015 CVPR
MFA [8]	50.8	78.7	87.7	47.9	75.1	84.6	41.6	70.5	80.6	30.0	57.2	69.7	2014 ECCV
kLFDA [8]	51.3	78.8	88.2	48.6	75.7	84.9	20.9	39.3	49.4	9.1	23.9	32.5	2014 ECCV
RBML [15]	54.0	80.7	88.6	51.3	75.9	84.1	35.6	63.0	74.3	29.8	55.3	67.0	2014 ECCV
LFDA [7]	54.1	79.4	87.8	49.9	75.0	83.8	28.9	50.9	62.1	30.5	55.1	66.9	2013 CVPR
RS-KISS [13]	54.6	79.5	88.1	49.5	74.4	83.6	33.4	61.1	72.4	29.1	52.9	65.4	2013 IEEE
KISSME [12]	47.1	75.5	85.6	41.8	69.4	79.7	33.9	58.9	69.3	25.9	53.6	66.7	2012 CVPR



Fig. 10. Example images from PRID 450S dataset [25].

that our proposed KRKISS method has the highest performance in all compared KISS-based methods. Meanwhile, it further proves the necessity of kernelized feature transform and the random subspace ensemble for KISS based methods.

Next, we compare our proposed KRKISS method with some state-of-the-art metric learning methods [6,7,8,12,13, 15,16,21]. The features we used in this experiment include LOMO [6], FFN [17], SCNCD [4], and ELF18 [19] which were reported with good performance on the PRID 450S dataset. The experimental results are shown in Table 7 and Fig 12. It can be seen that the KRKISS method achieves the best performance with three features (LOMO, SCNCD, ELF18) and the second best with FNN feature. The results of comparing with other state-of-the-art metric learning methods

Table 8

Comparison of state-of-the-art results reported with the PRID 450S database (P = 225). The cumulative matching scores (%) at rank 1, 5, 10, and 20 are listed. The best and the second best scores are respectively shown in red and blue.

Method	<i>r</i> = 1	<i>r</i> = 5	<i>r</i> = 10	<i>r</i> = 20	Reference
KRKISS	70.4	89.0	94.1	97.8	Ours
GOG [29]	66.4	87.6	93.8	97.5	2016 CVPR
NFST [21]	68.0	87.0	92.6	96.4	2016 CVPR
MLAPG [16]	55.5	80.1	88.3	94.7	2015 ICCV
LOMO + XQDA [6]	55.9	80.6	88.0	93.4	2015 CVPR
RBML [15]	54.0	80.7	88.6	93.9	2014 ECCV
RS-KISS [13]	54.6	79.5	88.1	94.0	2013 IEEE
KISSME [12]	46.4	74.9	84.9	92.3	2012 CVPR

demonstrate the effectiveness and the robustness of our proposed KRKISS algorithm.

4.2.2. Comparison to the state of the art

Next we compare our proposed method with the state-of-theart methods on the PRID 450S dataset. The compared methods and the experimental setups are same as to the experiments on the VIPeR dataset. The cumulative matching rates at rank 1, 5, 10, and 20 are listed in Table 8 and the CMC curves are shown in Fig 13. As we can see in Table 8, the rank-1, rank-5, rank-10 and rank-20 matching rates of our proposed method achieve 70.4%, 89.0%, 94.1% and 97.7%, respectively, which are the highest in all compared methods. The PRID 450S dataset is regarded as more realistic for person re-identification, and thus the experimental results on the PRID 450S dataset demonstrate that our proposed method is robust to different environments of person *Re*-ID.



Fig. 11. CMC curves on the PRID 450S dataset (P = 225). (a) Comparison of KISS based metric learning methods by using the LOMO feature. (b) Comparison of KISS based metric learning methods by using the FFN feature.



Fig. 12. CMC curves on the PRID 450S dataset (P = 225). (a) Comparison of different metric learning methods by using the LOMO feature. (b) Comparison of different metric learning methods by using the FFN feature. (c) Comparison of different metric learning methods by using the ELF18 feature.



Fig. 13. The CMC curves and rank-1 matching rates on the PRID 450S dataset.

4.3. Experiments on the CUHK01 dataset

CUHK01 [27] is another challenging person re-identification dataset which contains 971 pedestrians. Each pedestrian has four images captured from two different camera views in a campus environment. Camera A captures the frontal view or back view of people while camera B captures the side view. Large inter-camera variations are observed in this dataset which makes person reidentification challenging. All images have been normalized to



Fig. 14. Example images from CUHK01 dataset [27].

 160×60 pixels. Fig 14 shows some example images from this dataset. Different with the experimental settings of VIPeR dataset and PRID 450S dataset, the expriments of CUHK01 dataset can be divided into CUHK01(M = 1) and CUHK01(M = 2) because each pedestrian has more than two images. The experimental setting of



Fig. 15. CMC curves on the CUHK01 dataset (P = 486). (a) Comparison of KISS based metric learning methods with the single shot setting by using the LOMO feature. (b) Comparison of KISS based metric learning methods with the multi shot setting by using the LOMO feature. (c) Comparison of KISS based metric learning methods with the single shot setting by using the FFN feature. (d) Comparison of KISS based metric learning methods with the multi shot setting by using the FFN feature.

Comparison of different metric learning methods on the CUHK01 database (P = 486). The cumulative matching scores (%) at rank 1, 5 and 10 are listed. The best and the second best scores are respectively shown in red and blue.

Feature/Method	M = 1						M = 2						
	LOMO($d = 26,960$) FFN($d = 4096$)		4096)	LOMO($d = 26,960$)				FFN(d =	Reference				
	r = 1	<i>r</i> = 5	<i>r</i> = 10	r = 1	<i>r</i> = 5	<i>r</i> = 10	r = 1	<i>r</i> = 5	<i>r</i> = 10	r = 1	<i>r</i> = 5	<i>r</i> = 10	
KRKISS	61.4	81.7	87.4	42.7	65.2	74.6	66.9	86.9	91.9	46.2	70.3	79.1	Ours
NFST [21]	60.0	80.8	86.7	39.4	60.6	69.0	65.0	85.0	89.9	43.4	66.5	74.4	2016 CVPR
MLAPG [16]	58.4	79.0	85.5	24.8	46.5	56.9	64.7	86.6	91.6	26.8	50.3	61.3	2015 ICCV
XQDA [6]	55.8	78.6	85.7	34.5	55.6	63.9	62.8	83.9	90.5	39.7	60.1	68.4	2015 CVPR
MFA [8]	58.7	81.2	88.1	39.6	64.2	73.9	63.8	85.7	91.9	44.7	70.6	79.9	2014 ECCV
kLFDA [8]	58.9	80.8	86.9	32.0	53.8	64.2	64.7	84.0	90.4	34.1	58.5	69.0	2014 ECCV
RBML [15]	55.1	77.8	84.7	42.1	63.7	72.3	63.7	83.5	89.8	46.1	68.5	76.9	2014 ECCV
LFDA [7]	55.6	77.9	85.1	39.3	63.6	73.1	62.9	83.5	89.7	44.8	68.7	77.7	2013 CVPR
RS-KISS [13]	54.2	76.7	83.3	42.2	65.2	74.3	60.6	81.8	88.0	45.9	70.2	77.9	2013 IEEE
KISSME [12]	52.6	75.2	82.5	39.8	63.1	72.1	58.2	81.7	88.8	45.5	68.6	77.3	2012 CVPR

CUHK01(M = 1) is the single shot setting and CUHK01(M = 2) is the multi shot setting, which are common to [46] and [6] respectively.

4.3.1. Comparison of metric learning algorithms

Same as to the experiments in PRID 450S dataset, we compare our proposed KRKISS method with other three KISS based methods (KISSME [12], RS-KISS [13] and RBML [15]) by using two features (LOMO [6] and the FFN [17]) respectively. The experimental matching rate curves are shown in Fig 15. Meanwhile, we have performed some experiments to compare the KRKISS method with some state-of-the-art metric learning methods. The experimental results are shown in Table 9 and Fig 16.

As we can see in Fig 15, our proposed KRKISS method achieves better matching rates than other three KISS based methods no matter the experimental setting is the single shot setting or the multi shot setting. In Table 9 and Fig 16, compared with other state-of-the-art metric learning methods, the KRKISS method achieves the best rank-1 rate when the experimental setting is the single shot setting, and it is among the top two when the experimental setting is the multi shot setting. All the experimental results indicates that the KRKISS metric learning method is not only suit for the single-shot



Fig. 16. CMC curves on the CUHK01 dataset (P = 486). (a) Comparison of different metric learning methods with the single shot setting by using the LOMO feature. (b) Comparison of different metric learning methods with the multi shot setting by using the LOMO feature. (c) Comparison of different metric learning methods with the single shot setting by using the FFN feature. (b) Comparison of different metric learning methods with the single shot setting by using the FFN feature.

Comparison of state-of-the-art results reported with the CUHK01 database (P = 486). The cumulative matching scores (%) at rank 1, 5, 10, and 20 are listed. The best and the second best scores are respectively shown in red and blue.

Method	Method CUHK01(M = 1)				CUHK01(N	Reference			
	<i>r</i> = 1	<i>r</i> = 5	<i>r</i> = 10	<i>r</i> = 20	r = 1	<i>r</i> = 5	<i>r</i> = 10	<i>r</i> = 20	
KRKISS	66.9	83.0	88.7	92.7	72.9	89.8	94.0	96.9	Ours
NFST [21]	60.0	80.8	86.7	91.9	65.0	85.0	89.9	94.4	2016 CVPR
GOG [29]	59.8	77.6	84.3	89.9	66.4	84.9	90.4	94.5	2016 CVPR
MLAPG [16]	58.4	79.0	85.5	91.6	64.7	86.6	91.6	95.4	2015 ICCV
LOMO + XQDA [6]	55.8	78.6	85.7	91.3	62.8	83.9	90.5	94.6	2015 CVPR
RBML [15]	55.1	77.8	84.7	90.5	63.6	83.5	89.8	94.8	2014 ECCV
RS-KISS [13]	54.2	76.7	83.3	89.8	60.6	81.8	88.0	93.1	2013 IEEE
KISSME [12]	52.6	75.2	82.5	89.4	58.2	81.7	88.8	94.0	2012 CVPR

cases in person re-identification, but also suit for the multi-shot cases.

4.3.2. Comparison to the state of the art

Next we compare our proposed methods (KRKISS) with some state-of-the-art methods which were reported with good performance on the CUHK01 dataset. The compared methods and the experimental setups are same as to the experiments on the VIPeR dataset. The experimental results are shown in Table 10 and Fig 17. It can be observed that our proposed method achieves the best matching rates on CUHK01 (M = 1) and CUHK01(M = 2), which are 6.9% and 7.9% better than the second at rank-1 respectively. The significantly improvement of performance demonstrates that our proposed KRKISS method is also suitable to multi shot scenes in the applications of person *Re*-ID.

5. Conclusions and future works

In this paper, we have presented a novel metric learning method for person re-identification called Kernelized Random KISS (KRKISS). The KRKISS method can augment Guassian distribution of data by transforming the original features into kernelized feature. By doing so, the differences between kernelized features pairs can better fit to Gaussian distribution, and thus they can be more suitable for the Guassian assumption based models. Meanwhile, the SSS problem in covariance matrix estimation based methods can now be solved by a random subspace ensemble method because the influence of SSS problem would decreases or disappears when the dimension of subspace is greatly less than number of samples. The experiments on three publicly datasets, VIPeR, PRID 450S, CUHK01, show the effectiveness and the robust-



Fig. 17. The CMC curves and rank-1 matching rates on the CUHK01 dataset. (a) Comparison of some state-of-the-art methods with the single shot setting. (b) Comparison of some state-of-the-art methods with the multi shot setting.

ness of KRKISS method, and demonstrate that the KRKISS method is better than the state-of-the-art metric learning methods in most cases. It would be interesting to see that the KRKISS method can apply to other cross-view matching problem, such as face recognition.

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